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(54) Title: NOVEL FAS PROTEIN AND METHODS OF USE THEREOF

(57) Abstract

The present invention provides a novel form of the Fas protein, DNA encoding the protein cells expressing the recombinant DNA and methods of use thereof.

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#### NOVEL FAS PROTEIN AND METHODS OF USE THEREOF

#### Field of the Invention

This invention relates generally to the field of apoptosis and specifically to a novel Fas protein.

#### 5 Background of the Invention

Apoptosis is a normal physiologic process that determines individual cell death and ultimate deletion of the cell from tissue. For review see, Apoptosis the Molecular Basis of Cell Death, Tomei and Cope, eds., Current Communications in Cell and Molecular Biology 3, Cold Spring Harbor Laboratory Press, NY, 1991. Apoptosis is a process of programmed cell death involved in a variety of normal and pathogenic biological events and

can be induced by a number of unrelated stimuli. Changes
in the biological regulation of apoptosis also occur
during aging and are responsible for many of the
conditions and diseases related to aging.

Recent studies of apoptosis have implied that a common metabolic pathway leading to cell death may be initiated by a wide variety of signals, including changes in hormone levels, serum growth factor deprivation, chemotherapeutic agents and ionizing radiation. Wyllie (1980) Nature, 284:555-556; Kanter et al. (1984) Biochem. Biophys. Res. Commun., 155:324-331; and Kruman et al. (1991) J. Cell. Physiol., 148:267-273. Agents that affect the biological control of apoptosis thus have therapeutic utility in a wide variety of conditions.

Fas, also known as APO-1, is a cell surface protein belonging to the tumor necrosis factor/nerve growth

30 factor receptor family, each of whose members have been shown to be capable of mediating apoptosis. The cloning of Fas is described in PCT publication No. WO 91/10448; and European Patent Application Publication

Number 0510691. Fas is a transmembrane (TM) protein of

35 34,971 deduced molecular weight and an apparent molecular

weight of about 45,000 which may be due to glycosylation. The mature Fas molecule consists of 319 amino acid residues of which 157 are extracellular, 17 constitute the TM domain and 145 are intracellular. A variety of cell types express Fas on their surface. Interestingly, Fas expression is increased in activated T-cells including CD4+ and CD8+ cells.

Certain antibodies specific for Fas have been shown to induce death of cells that express Fas on their 10 surfaces, by an apoptotic mechanism. Early studies indicated that therapeutic uses of antibodies specific to Fas would be effective in treating a variety of diseases. Itoh et al. (1991) Cell, 66:233-243; Krammer et al. (1991) in Apoptosis: The Molecular Basis of Cell Death, 15 (Tomei and Cope, eds.), Cold Spring Harbor Laboratory Press, NY; Oehm et al. (1992) J. Biol. Chem., 267:10709-10715; and Rouvier et al. (1993) J. Exp. Med., 177:195-It has now been found that administration of anti-Fas antibodies can be lethal. Ogasawara et al. (1993) Nature, 364:806-809. It has also been found that 20 purified Fas blocks the cytocidal effects of anti-Fas. Oehm et al. (1992).

Increased levels of T cell surface Fas have also been associated with tumor cells and HIV-infected cells.

25 HIV-infected cells are more sensitive to anti-Fas antibodies, yet the significance of the association of Fas with HIV infection has not yet been determined.

The endogenous Fas ligand, responsible for recognizing Fas and inducing apoptosis, has not been identified, although some AIDS patients have been shown to have increased levels of anti-Fas autoantibodies. Oehm et al. (1991). Moreover, T cell mediated cytotoxicity has been shown to be involved in Fasmediated apoptosis.

All references cited herein both <u>infra</u> and <u>supra</u> are hereby incorporated herein by reference.

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#### Summary of the Invention

A novel native form of the Fas protein,
(hereinafter, FasΔTM) lacking the transmembrane region is
provided. DNA encoding FasΔTM and recombinant cells
expressing the DNA are also provided. Diagnostic and
therapeutic methods utilizing FasΔTM are also provided.

#### Brief Description of the Drawings

Figure 1 is a schematic diagram depicting the Fas genomic DNA structure around the transmembrane (TM) region.

Figure 2 is a schematic diagram depicting alternate splicing of the Fas RNA to produce Fas and Fas $\Delta$ TM mRNA.

Figure 3 depicts the nucleotide and amino acid residue sequences of Fas $\Delta$ TM.

Figure 4 depicts synthetic peptides used to raise antibodies useful in the detection of biologically important Fas molecules.

Figure 5 depicts survival of various cell lines transformed with vectors expressing Fas $\Delta$ TM or control, untransformed cell lines on exposure to Fas antibodies.

Figure 6 depicts survival of various cell lines transformed with vectors expressing Fas $\Delta$ TM or control, untransformed cell lines on exposure to Fas antibodies.

Figure 7 depicts the amount of Fas on the surface of various cell lines transformed with vectors expressing Fas $\Delta$ TM or control, untransformed cell lines.

Figure 8 depicts the activity of recombinant Fas $\Delta$ TM in preventing anti-Fas antibody induced cell death.

#### Detailed Description of the Invention

The present invention is to an isolated novel, secreted form of Fas protein, hereinafter designated FasΔTM and methods of use of FasΔTM. The invention further includes the cloned DNA encoding FasΔTM and recombinant cells expressing the DNA. The nucleotide and

amino acid residue sequences of Fas $\Delta$ TM are shown in Figure 3.

The genomic structure of the Fas gene, in the appropriate region, is depicted in Figure 1. 5 locations of the introns and exons are related to the different regions of Fas. The cloning of Fas∆TM is described in detail in the Examples below.

Figure 2 depicts the alternate splicing of the mRNA thought to result in the alternate forms of Fas however, 10 the invention is not limited by the mechanism by which Fas $\Delta$ TM is produced. Native Fas $\Delta$ TM lacks twenty-one amino acid residues including the TM region.

The invention includes other recombinant variations of Fas ATM which lack a portion of the TM region 15 sufficient to produce non-membrane bound protein. Preferably, the protein is secreted from the cell. term Fas $\Delta$ TM encompasses all the non-membrane-bound forms of the molecule lacking TM region amino acid residues. Figure 3 depicts the nucleotide and amino acid residue sequences of native Fas ATM.

One embodiment of the present invention is the DNA encoding Fas $\Delta$ TM. The DNA encoding Fas $\Delta$ TM includes, but is not limited to, the cDNA, genome-derived DNA and synthetic or semi-synthetic DNA. The nucleotide sequence of the cDNA encoding Fas $\Delta$ TM is shown in Figure 3. DNA includes modifications such as deletions, substitutions and additions, particularly in the noncoding regions. Such changes are useful to facilitate cloning and modify gene expression. Various substitutions can be made within the coding region that 30 either do not alter the amino acid residues encoded or result in conservatively substituted amino acid residues. Nucleotide substitutions that do not alter the amino acid residues encoded are useful for optimizing gene 35 expression in different systems. Suitable substitutions are known to those of skill in the art and are made, for

instance, to reflect preferred codon usage in the

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particular expression systems. Suitable conservative amino acid residue substitutions are known in the art and are discussed below.

Techniques for nucleic acid manipulation useful for
the practice of the present invention are described in a
variety of references, including, but not limited to,

Molecular Cloning: A Laboratory Manual, 2nd ed.,
Vol. 1-3, eds. Sambrook et al. Cold Spring Harbor
Laboratory Press (1989); and Current Protocols in
Molecular Biology, eds. Ausubel et al., Greene Publishing

Molecular Biology, eds. Ausubel et al., Greene Publishing and Wiley-Interscience: New York (1987) and periodic updates.

The invention further embodies a variety of DNA vectors having cloned therein the nucleotide sequence encoding Fas $\Delta$ TM. Suitable vectors include any known in 15 the art including, but not limited to, those for use in bacterial, mammalian and insect expression systems. Specific vectors are known in the art and need not be described in detail herein. The vectors may also provide 20 inducible promoters for expression of Fas $\Delta$ TM. Inducible promoters are those which do not allow constitutive expression of the gene but rather, permit expression only under certain circumstances. Such promoters may be induced by a variety of stimuli including, but not limited to, exposure of a cell containing the vector to a 25 ligand, chemical or change in temperature.

These promoters may also be cell-specific, that is, inducible only in a particular cell type and often only during a specific period of time. The promoter may further be cell cycle specific, that is, induced or inducible only during a particular stage in the cell cycle. The promoter may be both cell type specific and cell cycle specific. Any inducible promoter known in the art is suitable for use in the present invention.

The invention further includes a variety of expression systems transfected with the vectors.

Suitable expression systems include, but are not limited

to, bacterial, mammalian and insect. Specific expression systems are known in the art and need not be described in detail herein. It has been found that the baculovirus expression system described below provides expression of biologically active FasΔTM. For expressing FasΔTM for therapeutic purposes however, mammalian expression systems, such as Chinese hamster ovary (CHO) cells, may be preferred to ensure proper post-translational modification.

- The invention encompasses cells removed from animals, including man, transfected with vectors encoding FasΔTM and reintroduced into the animal. Suitable transfected cells also include those removed from a patient, transfected and reintroduced into the patient.
- Any cells can be treated in this manner. Suitable cells include, but are not limited to, cardiomyocytes and lymphocytes. For instance, lymphocytes, removed, transfected with the recombinant DNA and reintroduced into an HIV-positive patient may increase the half-life of the reintroduced T cells.

Preferably, for treatment of HIV-infected patients, the white blood cells are removed and sorted to yield the CD4 $^+$  cells. The CD4 $^+$  cells are then transfected with a vector encoding Fas $\Delta$ TM and reintroduced into the patient.

- Alternatively, the unsorted lymphocytes can be transfected with a recombinant vector encoding the FasΔTM gene under the control of a cell-specific promoter such that only CD4+ cells express the FasΔTM gene. In this case, an ideal promoter would be the CD4 promoter,
- $^{30}$  however, any suitable CD4 $^{+}$  T cell-specific promoter can be used.

Further, the invention encompasses cells transfected in vivo by the vectors. Suitable methods of in vivo transfection are known in the art and include, but are not limited to, that described by Zhu et al. (1993) Science, 261:209-211. In the case of in vivo

transfection, it is preferred that the transfection is cell-type specific or that the promoter is cell-specific.

Transgenic animals containing the recombinant DNA vectors are also encompassed by the invention. Methods of making transgenic animals are known in the art and need not be described in detail herein. For a review of methods used to make transgenic animals, see PCT publication no. WO 93/04169. Preferably, such animals express a recombinant FasATM gene under control of a cell-specific and, even more preferably, a cell cycle specific promoter.

Purification or isolation of Fas  $\Delta$ TM expressed either by the recombinant DNA or from biological sources such as sera can be accomplished by any method known in the art.

- 15 Since native and most recombinant Fas∆TM are secreted from the cells, purification is simplified by the fact that it appears in the supernatant of <u>in vitro</u> cultures and in sera <u>in vivo</u> and no cell disruption is required as is the case with Fas. Protein purification methods are
- known in the art. Generally, substantially purified proteins are those which are free of other, contaminating cellular substances, particularly proteins. Preferably, FasΔTM is more than eighty percent pure and most preferably FasΔTM is more than ninety-five percent pure.
- For clinical use as described below, FasΔTM is preferably highly purified, at least about ninety-nine percent pure, free of pyrogens and other contaminants.

Suitable methods of purification are known in the art and include, but are not limited to, affinity

30 chromatography, immunoaffinity chromatography, size exclusion chromatography, HPLC and FPLC. Any purification scheme that does not result in substantial degradation of the protein is suitable for use in the present invention.

The invention also includes the substantially purified Fas $\Delta$ TM protein having the amino acid residue sequence depicted in Figure 3 and any protein lacking a

sufficient portion of the TM region to be secreted from The invention encompasses functionally equivalent variants of Fas ATM which do not significantly affect its properties. For instance, conservative substitutions of amino acid residues, one or a few amino acid deletions or additions, and substitution of amino acid residues by amino acid analogs are within the scope of the invention.

Amino acid residues which can be conservatively 10 substituted for one another include but are not limited glycine/alanine; valine/isoleucine/leucine; asparagine/glutamine; aspartic acid/glutamic acid; serine/threonine; lysine/arginine; and phenylalanine/tyrosine. Any conservative amino acid substitution which does not significantly affect the 15 properties of Fas $\Delta$ TM is encompassed by the present invention.

mRNA encoding FasΔTM has been detected in a variety of human organs and tissues. These include liver; heart; peripheral blood lymphocytes (PBLs), both activated and normal; placenta; fibroblasts, both normal and phorbol ester treated and SV40 infected; and several cell lines including U937, WIL-2 and IM9. Fas∆TM has also been found to be secreted from the cell rather than remain 25 membrane-bound, even though it retains the cytoplasmic region of the membrane-bound form of the protein. can thus be detected in sera as a soluble protein. antibody that recognizes Fas is suitable for use in recognizing FasΔTM.

30 In another embodiment, diagnostic methods are provided to detect the expression of Fas $\Delta$ TM either at the protein level or the mRNA level. The soluble  $Fas\Delta TM$ protein is likely to be found in the sera of patients with diseases associated with apoptosis defects, and is 35 therefore useful as a diagnostic tool for detecting and monitoring biological conditions associated with such apoptosis defects.

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Fas∆TM can be detected by any antibody, either polyclonal or monoclonal, that recognizes Fas. distinction between Fas and Fas ATM can be determined by the solubility of the protein. Fas is membrane bound and 5 can thus be removed from soluble proteins by removal of cells and/or membranes. FasΔTM remains soluble. Alternatively, antibodies specific for FasΔTM and not Fas are encompassed by the present invention. antibodies can be generated by using  $Fas\Delta TM$  as the 10 antigen or, preferably, peptides encompassing the region in Fas $\Delta$ TM that differs from Fas, the TM region. Examples of such peptides are depicted in Figure 5. detecting proteins using antibodies and of generating antibodies using proteins or synthetic peptides are known in the art and need not be described in detail herein.

Fas $\Delta$ TM protein expression can also be monitored by measuring the level of mRNA encoding Fas- $\Delta$ TM. for detecting specific mRNA species is suitable for use in this method. This is easily accomplished using the polymerase chain reaction (PCR). Preferably, the primers chosen for PCR flank the TM region so as to provide a product that is measurably distinct in size between Fas and Fas $\Delta$ TM. Alternatively, Northern blots can be utilized to detect the specific mRNA species either by size or by probes specific to the mRNA encoding the TM region.

The invention also encompasses therapeutic methods and compositions involving treatment with Fas∆TM. native or recombinant Fas $\Delta$ TM is suitable for use in this 30 composition. Both should be substantially pure and free of pyrogens. It is preferred that the recombinant FasATM be produced in a mammalian cell line so as to ensure proper glycosylation. Fas∆TM may also be produced in an insect cell line and will be glycosylated.

35 For therapeutic compositions, a therapeutically effective amount of substantially pure Fas∆TM is suspended in a physiologically accepted buffer including, but not limited to, saline and phosphate buffered saline (PBS) and administered to the patient. Preferably administration is intravenous. Other methods of administration include but are not limited to,

5 subcutaneous, intraperitoneal, gastrointestinal and directly to a specific organ, such as intracardiac, for instance, to treat cell death related to myocardial infarction.

Suitable buffers and methods of administration are known in the art. The effective concentration of FasΔTM will need to be determined empirically and will depend on the type and severity of the disease, disease progression and health of the patient. Such determinations are within the skill of one in the art. Moreover, FasΔTM is a human protein normally found in the sera; administration of exogenous human FasΔTM is not likely to induce reactions such as anaphylactic shock or the production of antibodies. The upper concentration of FasΔTM for therapeutic use is thus not limited by these physiological considerations.

Administration of FasATM results in an increased extracellular concentration of FasATM which competitively binds the Fas ligand and therefore prevents or ameliorates apoptotic signals transmitted by Fas to the cell. The therapeutic method thus includes, but is not limited to, inhibiting Fas-mediated cell death. For instance, tumor necrosis factor (TNF) and Fas-specific antibodies are known to induce apoptosis and even whole animal death by binding to Fas. Inhibition of this interaction of Fas and ligands which induce it to trigger apoptosis thus will reduce apoptosis.

Suitable indications for therapeutic use of Fas ATM are those involving Fas-mediated cell death and include, but are not limited to, conditions in which there is inappropriate expression or up-regulation of Fas or the Fas ligand. Such indications include, but are not limited to, HIV infection, autoimmune diseases,

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cardiomyopathies, neuronal disorders, hepatitis and other liver diseases, osteoporosis, and shock syndromes, including, but not limited to, septicemia.

Methods of treatment with Fas∆TM also include other 5 mechanisms of increasing the extracellular concentration of Fas $\Delta$ TM. These include, but are not limited to, increasing cellular expression of FasATM. methods of increasing cellular expression of FasΔTM include, but are not limited to, increasing endogenous 10 expression and transfecting the cells with vectors encoding Fas∆TM. Cellular transfection is discussed above and is known in the art. Increasing endogenous expression of Fas $\Delta$ TM can be accomplished by exposing the cells to biological modifiers that directly or indirectly increase levels of Fas $\Delta$ TM either by increasing expression 15 or differential processing of the Fas $\Delta$ TM over Fas or by decreasing Fas ATM degradation. Suitable biological modifiers can be determined by exposing cells expressing  ${\tt Fas}\Delta{\tt TM}$  under the control of the native  ${\tt Fas}$  promoter to 20 potential biological modifiers and monitoring expression Expression of Fas $\Delta$ TM can be monitored as of Fas $\Delta$ TM. described above either by protein or mRNA levels. Biological modifiers can be any therapeutic agent or chemical known in the art. Preferably, suitable 25 biological modifiers are those lacking substantial cytotoxicity and carcinogenicity.

Likewise, biological modifiers which reduce endogenous levels of FasATM are encompassed by the invention as is a method of increasing Fas-mediated cell death by decreasing endogenous levels of FasATM. The method of determining suitable biological modifiers is as discussed above, except that the endpoint is decreased levels of FasATM. Other methods of decreasing endogenous levels of FasATM include, but are not limited to, antisense nucleotide therapy and exposure to anti-FASATM antibody. Both these methods are known in the art and their application will be apparent to one of skill in the

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art. Suitable indications for decreasing endogenous levels of Fas∆TM will be any which Fas-mediated cell death is appropriate. These include, but are not limited to, various types of malignancies and other disorders resulting in uncontrolled cell growth such as eczema.

The following examples are provided to illustrate but not limit the present invention.

#### Example 1

#### Cloning of Fas-ΔTM

- Nucleic acid sequences encoding FasΔTM were cloned as follows. Unless otherwise specified, all cloning techniques were essentially as described by Sambrook et al. (1989) and all reagents were used according to the manufacturer's instructions.
- mRNA was obtained from human lymphocytes and PCR was used to make cDNA specific for the Fas $\Delta$ TM mRNA. The lymphocytes were obtained and processed as follows. 35 ml of blood was obtained by venipuncture from a normal 42 year old male and immediately added to 350  $\mu$ l 15% EDTA.
- 35 ml PBS was added to the blood and 17 ml of the blood suspension was layered on 12.5 ml Ficoll Paque (Pharmacia). This was then centrifuged at 1,800 rpm for 30 minutes in a swinging bucket rotor and a DynacII centrifuge.
- The plasma was aspirated and the lymphocyte layer collected and added to two volumes of PBS containing 0.9 mM Ca<sup>2+</sup> and 0.5 mM Mg<sup>2+</sup> (PBS/Ca/Mg buffer). The cells were washed once with the PBS/Ca/Mg buffer and resuspended in RPMI (Gibco/BRL) medium containing 2 g/l glucose and 10% fetal bovine serum (FBS) at 2 x 10<sup>6</sup> cells/ml. The cell yield was 3.5 x 10<sup>7</sup> with greater than 98% viability. Concanavalin A (Sigma) was added to a final concentration of 30 μg/ml and the cells were incubated at 37°C for 72 hr.
- 35 The cells were then processed for RNA isolation as follows. Total RNA was isolated from 3  $\times$  10<sup>6</sup> cells using

the guanidinium thiocyanate single-step RNA isolation method according to "Current Protocols in Molecular Biology" (1991). The first strand cDNA used in the PCR reaction was synthesized from 6.4 μg of total RNA and resuspended in 100 μl water, according to the method described by Zapf et al. (1990) J. Biol. Chem., 265:14892-14898. The FasΔTM cDNA was synthesized using PCR. The forward primer:

5'-GATTGCTTCTAGACCATGCTGGGCATCTGGACCCTCCTACC-3' contained 10 an XbaI restriction site, and encoded the initiation methionine codon and first eight codons of Fas. The reverse primer:

The PCR reaction was performed by adding to 1  $\mu$ l of template cDNA (as described above, diluted 1:100), 100 pmoles of each primer, 2.5 units Amplitaq, 76.5  $\mu$ l H<sub>2</sub>O and 10  $\mu$ l buffer. The reaction proceeded at 94°C for 1 min,

- 55°C for 2 min, 72°C for 3 min for 35 cycles, 72°C for 7 min and was stored at 4°C. 15 μl of the reaction mix was loaded on a 1% agarose gel and a band of the appropriate molecular weight was detected. The remaining reaction mix was extracted with phenol/chloroform, ethanol
  25 precipitated and resuspended in 80 μl Trice FDTA (ME)
- precipitated and resuspended in 80  $\mu$ l Tris-EDTA (TE). To the 80  $\mu$ l sample the following was added: 5  $\mu$ l SalI; 5  $\mu$ l XbaI; 10  $\mu$ l SalI reaction buffer. In another reaction, 60  $\mu$ l H<sub>2</sub>O was added to 20  $\mu$ l pBluescript at 0.65 mg/ml, 5  $\mu$ l SalI, 5  $\mu$ l XbaI and 10  $\mu$ l SalI reaction
- buffer. Both tubes were incubated at 37°C for 2 hours and the reactions were run on a 1% preparative agarose gel. Bands corresponding to the digested DNA were excised from the gel and purified by Elutip® according to the manufacturer's instructions. The purified FasΔTM
- 35 cDNA was resuspended in 20  $\mu l$  TE buffer and the pBluescript in 40  $\mu l$  TE buffer.

The DNA samples were ligated in a reaction mixture containing 2  $\mu$ l vector, 8  $\mu$ l Fas $\Delta$ TM cDNA, 2  $\mu$ l 10 mM ATP, 2  $\mu$ l 10 x ligation buffer, 2  $\mu$ l T4 DNA ligase (New England Biolabs) and 4  $\mu$ l H<sub>2</sub>O. The control reaction contained no Fas $\Delta$ TM DNA. After allowing the ligation to continue for 6 hr at 14°C the DNA was used to transform  $DH5\alpha$  cells (Gibco) according to the manufacturer's instructions. Briefly, 200  $\mu l$  cells were added to the ligation mix and kept on ice for 45 min. The cells were 10 heat shocked for 90 sec at 42°C and then placed on ice. 3 ml L broth was then added and the cells were incubated for 1 hr at 37°C and plated on L broth agar plates containing 100 mM ampicillin, 20  $\mu$ l 4% X-Gal and 50  $\mu$ l 100 mM isopropyl-1- $\beta$ -D-thiogalactoside (IPTG). The cells 15 were allowed to form colonies by incubating overnight at 37°C. Positive colonies were grown overnight in L broth plus ampicillin, the plasmids were obtained by an alkaline lysis procedure and digested with SalI and XbaI according to the manufacturer's instructions. plasmid containing the appropriate insert was prepared on 20 a large-scale and the insert was sequenced by the dideoxy The sequence obtained is presented in Figure 3. The plasmid containing the recombinant cDNA encoding Fas $\Delta$ TM was designated pBluescript-Fas $\Delta$ TM.

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#### Example 2

#### Analysis of Fas Genomic Structure

The intron-exon organization in the Fas TM region was determined by PCR. Primers were designed to flank each of the putative introns, 1 and 2 (see Figure 2).

The forward and reverse primers flanking intron 1 were 5'-GATTGCTTCTAGAGGAATCA TCAAGGAATGCACACTC-3' and 5'-GTTGTTTGTCGACC CAAACAATTAGTGGAATTGGCAA-3' respectively, and the forward and reverse primers for intron 2 were 5'-AGATCTGCGGCCGCAT

TGGGGTGGCTTTGTCTTCTT-3' and 5'-GTTGTTTGTCGACGTTTTCCT

TTCTGTGCTTTCTGCA -3' respectively. XbaI and SalI

restriction enzyme sites and NotI and SalI restriction enzyme sites were included at the 5' ends of the intron 1 and 2 primers respectively to facilitate cloning of the PCR products. PCR was performed according to the 5 manufacturer's instructions (Perkin Elmer Cetus) using human genomic DNA (Clontech) (5  $\mu$ g) as template. cycles of PCR were performed in a Perkin Elmer Cetus DNA Thermal Cycler with each cycle consisting of a 94°C, 1 min denaturation step, a 55°C, 2 min annealing step, and 10 a 72°C, 3 min extension step. An additional 7 minute extension step was included after the last cycle. PCR products were then incubated with 5 units of DNA polymerase I, Klenow fragment at 37°C, 30 min, extracted with phenol/chloroform/isoamylalcohol (1:1:0.04) followed 15 by chloroform/isoamylalcohol (24:1) and recovered by ethanol precipitation.

The intron 1 and intron 2 PCR products were digested with XbaI and SalI and NotI and SalI respectively, agarose gel purified, ligated into pBluescript SK(-) and introduced into E. coli strain HB101 by the methods described in Example 1. Plasmid DNA was isolated using a Promega Magic miniprep kit according to the method described by the manufacturer. Plasmid DNA was sequenced directly by the dideoxy chain termination method using the Sequenase Version 2.0 DNA Sequencing kit according to the manufacturer's instructions (USB). The sequence obtained is depicted in Figure 3.

#### Example 3

#### Expression of Recombinant Fas-ΔTM

- In order to express recombinant FasΔTM in the baculovirus system, the plasmid obtained in Example 1 was used to generate a second FasΔTM vector, designated pBlueBacIII-FasΔTM, by a PCR methodology as described in Example 1. The forward primer,
- 35 5'-TTTCCCGGATCCACAACCATGCTGGGCATCTGGACCCTCCTA-3' contained the convenient <a href="mailto:BamHI">BamHI</a> restriction site, and a

Kozak consensus sequence ACAACC immediately preceding the initiation codon, and encoded the first seven amino acids The reverse primer 5'-CCCCATGGCTAGACCAAGCTTTG GATTTCATT-3' encoded a termination codon, a NcoI site, 5 and the last seven amino acids of Fas. Five recombinant plasmids were isolated. Two of them were sequenced by the dideoxy terminator method (Sanger et al. 1977) using sequencing kits according to the manufacturer's instructions (USB, Sequenase version 2.0). sequenced using internal primers.

Clone 3, which did not contain any PCR errors in the sequence, was used to generate recombinant viruses by in vivo homologous recombination between the overlapping sequences pBlueBac III-FASATM-3 and AcNPV wild type 15 baculovirus. After 48 hours post-transfection in insect Spodoptera frugiperda clone 9 (SF9) cells, the recombinant viruses were collected, identified by PCR and further purified. Standard protocols for plasmid cloning were employed (Maniatis et al. 1982). Standard procedures for selection, screening and propagation of 20 recombinant baculovirus were performed according to the manufacturer's instructions (Invitrogen). After 48 hours post-transfection, the recombinant viruses were collected and purified. The molecular mass, on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-25 PAGE), of the protein produced in the baculovirus system is identical to the predicted molecular mass of Fas $\Delta$ TM according to the amino-acid sequence and the recombinant protein was also recognized by an anti-Fas antibody

#### Example 4

#### Expression of Fas ATM in Mammalian Systems

The Fas $\Delta$ TM coding sequence was excised from pBluescript-FasΔTM with XbaI and SalI, and introduced 35 into plasmids pCEP7 and pREP7 (Invitrogen) at compatible NheI and XhoI sites to generate clones Fas∆TM-1 and

(Medical and Biological Laboratories, Nagoya, Japan).

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Compatibility of the control of

FasΔTM-7, respectively. pCEP7 was generated by removing the RSV 3'-LTR of pREP7 with XbaI/Asp718, and substituting the CMV promoter from pCEP4 (Invitrogen). To generate a compatible XbaI site, pCEP4 was first cleaved with SalI, and ligated to an oligonucleotide adapter containing an external SalI site, and an internal NheI site. pCEP4 was then cleaved with NheI and Asp718 and the purified CMV promoter was ligated into pREP7 to generate pCEP7. 25 μg of each FasΔTM-containing plasmid was electroporated into the B lymphoblastoid cell line WIL-2, and stable hygromycin resistant transformants were selected.

#### Example 5

# Anti-Fas Antibody Induced Death of the 15 Wild Type B Lymphoblastoid Cell Line WI-L2-729 HF2 and the Wild Type Cell Transformed by FasΔTM-1 and FasΔTM-7

 $2x10^5$  WIL-2, Fas $\Delta TM-1$  and Fas $\Delta TM-7$  transformed WIL-2 cells were grown in RPMI supplemented with 10% fetal 20 bovine serum (FBS). After washing with fresh medium, the cells were suspended in RPMI supplemented with 10% FBS, 50 ng anti-Fas antibody was added, and the kinetics of cell death were analyzed by flow cytometry with FACScan. This method is based on the measurement of cells which shrink and are permeable to propidium iodide (PI) following their death. There was no difference in survival of all three cell lines in the control, but upon addition of anti-Fas antibodies, cells transformed by Fas $\Delta$ TM were less sensitive: by 26 hrs treatment 30 approximately 50%, 40%, and 16% of wild type WIL-2, Fas $\Delta$ TM-7, and Fas $\Delta$ TM-1 transformants died respectively (Fig. 5).

In another series of experiments, cells were initially grown for 24 hrs and then anti-Fas antibodies were added under the assumption that Fas∆TM was secreted and should accumulate in medium. In this case the

sensitivity of transformed cell lines was lower then without preincubation: after 26 hrs treatment with antibodies about 45%, 16%, and 5% of wild type WIL-2, Fas $\Delta$ TM-7, and Fas $\Delta$ TM-1 transformants died respectively (Fig. 6).

To ensure that lower sensitivity of transformants is not caused by down regulation of Fas expression the amount of Fas on the surface of all three cell lines was compared. The cells were treated by standard procedures 10 with monoclonal mouse anti-Fas antibodies, biotin-labeled anti-mouse IgM antibodies, and finally stained with FITClabeled streptavidin. Analysis of stained cells using FACScan showed that there were no differences in the amount of Fas on the surface of wild type WIL-2 and the 15 transformants (Fig. 7). Thus WIL-2 transformed with Fas $\Delta$ TM are less sensitive to the cytotoxic effect of anti-Fas antibodies. This effect may be explained at least in part by secretion of Fas $\Delta$ TM by transformants. Thus, increased cellular expression of Fas $\Delta$ TM results in 20 inhibition of Fas-mediated cell death.

#### Example 6

# Recombinant FasΔTM Prevents Cell Death Induced by Anti-Fas Antibodies

in a baculovirus system, insect cells transfected with wild type baculovirus and recombinant containing FasATM, as described in Example 3, were homogenized in water or in buffer containing 0.05% nonionic detergent Triton X-100 and centrifuged. Anti-Fas antibodies were preincubated 2 hrs at room temperature with aliquots of soluble and insoluble fractions, added to WIL-2, and cell death was analyzed after 24 hrs by flow cytometry as described above. There was no effect on cell viability of both soluble and insoluble fractions from insect cells transfected with the wild type baculovirus. At the same time soluble and insoluble fractions from insect cells

transfected with baculovirus recombinant containing
Fas-ΔTM inhibited death of WIL-2 induced by anti-Fas
antibodies. Activity of the insoluble fraction was
approximately ten times higher than that of the soluble
fraction (Fig. 8). Thus, recombinant FasΔTM can compete
with Fas on the cell surface for binding antibodies and
preventing Fas-mediated cell death.

#### Example 7

#### FasΔTM transcript analysis by RT-PCR

- 10 Native Fas ATM transcripts were identified by RT-PCR and acrylamide gel electrophoresis. PCR primers were designed around the Fas TM region so that the Fas and FasΔTM transcripts would yield 296 bp and 233 bp PCR products, respectively. The forward primer was 15 5'-GACCCAGAATACCAAGTGCAGATGTA-3' and the reverse primer was 5'-CTGTTTCAGGATTTAAGGTTGGAGATT-3'. cDNA was synthesized from poly(A)+ or total RNA isolated from various human tissues and cell lines. These included heart, liver, activated and non-activated peripheral 20 blood lymphocytes (PBLs), placenta and fibroblast cell lines. PCR was performed as described in Example 1 using the cDNA as templates (10-100 ng/ml) and products were analyzed on 7% acrylamide/TBE gels.
- All tissues and cell lines tested contained Fas and FasΔTM transcripts by this analysis. This suggests that cell death in these tissues can be modulated by the amounts (and ratios) of Fas and FasΔTM. Interestingly, liver contained the largest amounts of FasΔTM transcripts suggesting FasΔTM may be secreted into serum.
- Although the foregoing invention has been described in some detail by way of illustration and example for purposes of clarity of understanding, it will be apparent to those skilled in the art that certain changes and modifications may be practiced. Therefore, the
- description and examples should not be construed as limiting the scope of the invention, which is delineated by the appended claims.

#### We claim:

- 1. A composition comprising a purified nucleotide sequence encoding FasΔTM.
- 2. The nucleotide sequence according to claim 1 5 wherein the sequence is cDNA.
  - 3. The nucleotide sequence according to claim 1, having the nucleotide sequence depicted in Figure 3.
  - 4. A recombinant DNA vector comprising the nucleotide sequence according to claim 1.
- 10 5. The recombinant DNA vector according to claim 4 wherein expression of the sequence encoding Fas $\Delta$ TM is under control of an inducible promoter.
- 6. The recombinant DNA vector according to claim 4, selected from the group consisting of pBlueBACIII- Fas $\Delta$ TM-3, pBluescript-Fas $\Delta$ TM, Fas $\Delta$ TM-1 and Fas $\Delta$ TM-7.
  - 7. A cell transfected with the recombinant DNA vector according to claim 4.
  - 8. A transgenic animal comprising the recombinant DNA vector according to claim 4.
- 9. A substantially purified protein comprising the amino acid sequence of Figure 4.
- 10. A substantially purified protein comprising the Fas amino acid sequence but lacking a portion of the transmembrane region sufficient to result in a secreted protein that is not membrane bound.

- 11. The protein according to claim 10 wherein the protein is secreted.
- 12. The protein according to claim 10 wherein the protein is expressed by recombinant DNA.
- 5 13. The protein according to claim 10 wherein the protein is native protein.
  - 14. An antibody which recognizes Fas $\Delta$ TM but not Fas.
- 15. The antibody according to claim 14 selected from the group consisting of polyclonal and monoclonal.
  - 16. A composition comprising substantially purified Fas  $\Delta$ TM and a physiologically acceptable buffer.
  - 17. A method of detecting the presence of Fas $\Delta$ TM in a biological sample comprising the steps of
    - a) obtaining the biological sample;
  - b) adding anti-Fas antibodies to the biological sample;
  - c) maintaining the biological sample under conditions that allow the anti-Fas antibodies to complex with Fas $\Delta$ TM; and
    - d) detecting the complexes formed.
    - 18. The method according to claim 17 wherein the biological sample is sera.
- 19. The method according to claim 17 wherein the 25 anti-Fas antibody is specific for Fas∆TM.
  - 20. A method for detecting the expression of DNA encoding Fas $\Delta$ TM in a biological sample comprising

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identifying the presence in the biological sample of mRNA encoding Fas $\Delta$ TM.

- 21. The method according to claim 20 wherein the method for identifying the Fas∆TM mRNA is polymerase
  5 chain reaction and the primers flank the mRNA encoding the transmembrane region.
  - 22. The method according to claim 20 wherein the method for identifying the Fas $\Delta$ TM mRNA is Northern blotting.
- 23. A method of treating Fas-mediated cell death in a patient comprising administering to a patient a therapeutically effective amount of FasΔTM.
- 24. A method of protecting cells from Fas-mediated cell death comprising the steps of increasing the
   15 endogenous concentration of FasΔTM.
  - $25.\,$  The method according to claim 24 wherein the Fas  $\Delta TM$  is exogenous.
  - 26. The method according to claim 24 wherein the Fas $\Delta$ TM is expressed by the cell.
- 20 27. The method according to claim 26 wherein the Fas $\Delta$ TM is expressed by a recombinant gene.
  - 28. The method according to claim 27 wherein expression of the gene is under the control of an inducible promoter.

## FAS GENOMIC DNA STRUCTURE (TM REGION)

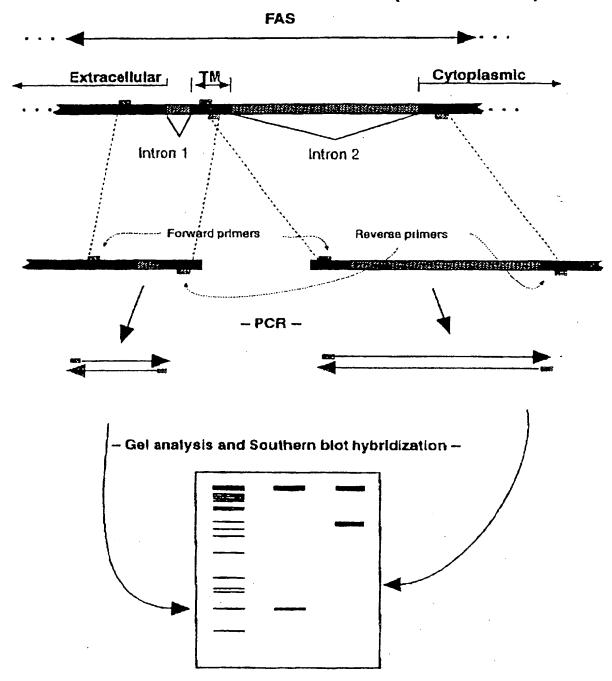


FIGURE [

#### FIGURE 2

## **SPLICING (TM REGION)**

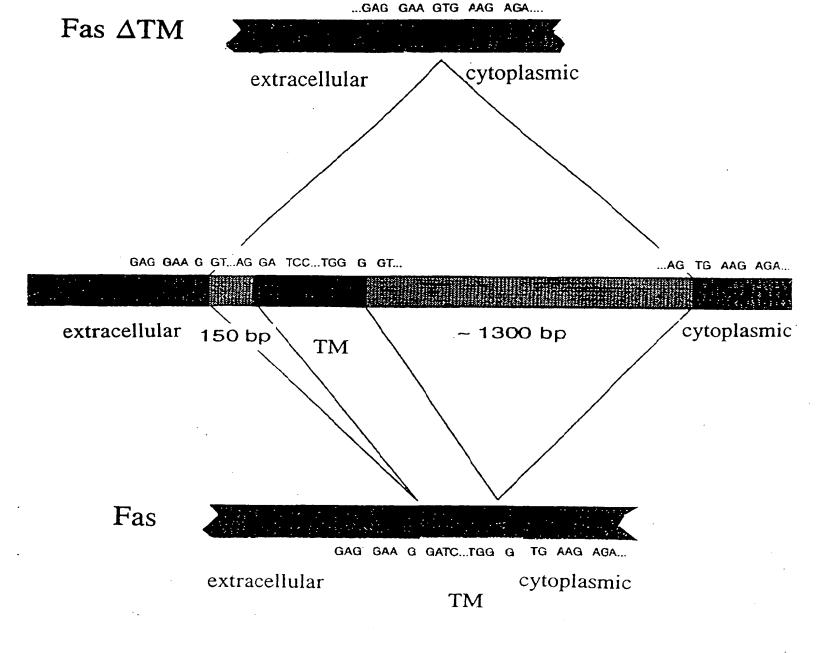


FIGURE 3-1

Nucleotide and Amino Acid Residue Sequences of Fas ATM

acacacctgaggccagccctggctgcccaggcggagctgcctcttctccgggggttggtggaccggctcagtacggagttggggaa GCTCTTTCACTTCGGAGGATTGCTCAACAACC

TCC TCG TTA AGA Arg +1 GCT Ala Val  ${\tt GTT}$ TCT Ser CTT ACG ' Leu Thr Val Leu CTG GTT ATC TGG ACC CTC CTA LLINE ILE Trp Thr Leu Leu Pro Leu GGC Gly Leu CTGATG Met

GTT Val ACT Thr GTT Val ACT Thr AAG Lys AGG Leu TTG GAA Glu 20 TTG Leu GGA Gly AAG Lys TCC GAC ATC AAC Asp Ile Asn CAA GTG ACT G Gln Val Thr 1 10 GCC Ala Asn AAT GTT Val

GGTG1yCCA Pro Pro CCT TGT Cys CCC Pro AAG Lys CAT His TGC Cys TTC CAA Gln GGC G1y 40 CAT GAT (His Asp Asp CAT His CTG Leu GGC Gly Glu GAA  ${
m TTG}$ Leu AAC Asn Gln CAG Thr ACT GAG

GGG Gly Glu GAA CAA Gln 70 TGC Cys CCC Pro GTG Val TGC Cys GAC Asp CCA Pro GAA Glu GAT 666 Gly AAT GTC Val ACA Thr TGC GAC Asp AGG GCTAla AAA Lys AGG Arg GAA Glu

GAA Glu GAT Cys  $\mathtt{TGT}$ Leu TTGAGA TGT Cys AGA Arg AGA Arg TGC Cys AAA Lys TCC Ser TCT Ser TTTPhe CAT His 80 GCC Ala AAA Lys GAC Asp TAC ACA Tyr Thr GAG Glu AAG Lys

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AAC Asn	ATC Ile	AAA Lys 160	ACA Thr	ACA Thr	AAG Lys
CCA	ATC	CAG	GAA	ATG	ATC
Pro	Ile	Gln	Glu	Met	Ile
AAA Lys	GGA Gly	GTA Val	CCT Pro 180	GTC Val	GAG Glu
TGT	CAT	GAA	AAT	GGA	GAT
Cys		Glu	Asn	Gly	Asp
AGA Arg	GAA Glu	AAG Lys	TTA Leu	GCT Ala 200	ATA Ile
TGC	TGT	aga	ACC	ATT	AAA
Cys	Cys	Arg	Thr	Ile	Lys
AAG Lys 110	AAA Lys	AAG Lys	CCA Pro	ACT Thr	GCC Ala 220
ACC	ACC	GTG	TCT	ACC	GAA
Thr	Thr	Val	Ser	Thr	Glu
AAT	TGC	GAA	GAA	ATC	AAT
Asn	Cys	Glu	Glu	Ile	Asn
CAG	CCT	GAG	CAT	TAT	GTC
Gln	Pro	Glu	His	Tyr	Val
ACC Thr	GAC Asp	AAA Lys 150	TCT Ser	AAA Lys	GGT Gly
CGG	TGT	TGC	GGT	AGT	AAT
Arg	Cys	Cys	Gly	Ser	Asn
ACC Thr	CAC His	AAG Lys	CAA Gln 170	TTG	AAG Lys
TGC	GAA	ACC	AAC	GAC	CGA
Cys	Glu	Thr	Asn	Asp	Arg
AAC Asn *	TGT Cys	AAC Asn	GAA Glu	GTT Val 190	GTT Val
ATA	GTA	AGC	AAG	GAT	TTT
Ile	Val	Ser	Lys	Asp	Phe
GAA Glu 100	ACT Thr	ACC Thr	AGA Arg	TCT Ser	GGC Gly 210
GTG	TCT	CTC	CAC	TTA	AAA
Val	Ser	Leu	His	Leu	Lys
GAA Glu	AAC Asn 120	ACA Thr	AAG Lys	AAT Asn	GTT Val
TTA	TGT	TGC	AGA	ATA	CAA
Leu	Cys	Cys		Ile	Gln
GGC Gly	TTT Phe	GAA Glu 140	TGC Cys	GCA Ala	AGT Ser
CAT	TTT	AAG	ACA	GTG	CTA
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GCC AAT C
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CTC AAA AAA Leu Lys Lys
CTC Leu
GAT ASP 260
AAA Lys
ACA TTG ATT AAA Thr Leu Ile Lys
TTG Leu
ACA Thr
GAC
TAT Tyr
GAA GCG Glu Ala
3 AAA C s Lys C
GGA AAG AAA Gly Lys Lys 250
66A 61y

AAT Asn
GA rrg
C TTC AGA AAT n Phe Arg Asn 290
C E E
A AA
TC? Se1
AAT Asn
AGT GAC TCA GAA AAT TCA AAC Ser Asp Ser Glu Asn Ser Asn
TCA Ser
GAC Asp
AGT Ser
AAG GAC ATT ACT Lys Asp Ile Thr 280
ATT Ile
GAC Asp 280
AAG Lys
ATC CTC AAG Ile Leu Lys
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CAG ACT ATC
CAG ACT Gln Thr
CA(
ATT Ile
AAA Lys
GAG AAA Glu Lys

GAA ATC CAA AGC TTG GTC TAG AGTGAAAAACAACTACAGTTCTGAGTATATGCAATTAGTGTTTGAAAAGATTCTTA Glu ile Gln Ser Leu Val End 298

ATAGCTGGCTGTAAATACTGCTTGGTTTTTACTGGGTACATTTTATCATTTATGCGCTGAAGAGCCAACATATTTGTAGATTTTT

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GTATGTTAGTACAAATGTCTATCCACAGGCTAACCCCCACTCTATGAATCAATAGAAGAAGCTATGACCTTTTGCTGAAATATCAGTTA

CTGAACAGGCCAGGTTTTGCCTCTAAATTACCTCTGATAATTCTAGAGATTTTACCATATTTCTAAACTTTGTTTATAACTCTGAG

FIGURE 3-4

angatcatattatatatatatatatttgagtgcagaatttaaataaa
CATATTATACAATATTTCAATTGTGAATTCACATAGAAAACATTAAATTTATAATGTTTGACTATTATATATGTGTGTATTTTACTG
GCTCAAAACTACCTACTTCTCAGGCATCAAAAGCATTTTGAGCAGGAGAGTATTACTAGAGCTTTGCCACCTCTCCATTTTTGC
CTTGGTGCTCATCTTAATGGCCTAATGCACCCCCAAACATGGAAATATCACCAAAAAATACTTAATAGTCCACCAAAAAGGCAAGACTG
CCCTTAGAAATTCTAGCCTGGTTTGGAGATACTACTGCTCTCAGAGAAAGTAGCTTTGTGACATGTCATGAACCCATGTTTGCAATC
AAAGATGATAAAATAGATTCTTATTTTCCCCCCCCCCGAAAATGTTCAATAATGTCCCATGTAAAACCTGCTACAAATGGCAGCTT
ATACATAGCAATGGTAAAATCATCTGGATTTAGGAATTGCTCTTGTCATACCCTCAAGTTTCTAAGATTTAAGATTCTCCTTACT
actatcctacgtttaaatatctttgaaagtttg <u>tattaa</u> atgtgaattttaagaaataatattttttttaggaaataat
AAGATAGTTATAAAĊTGAAGCAGATACCTGGAACCCTAAAGAACTTCCATTTATGGAGGATTTTTTGCCCCCTTGTGTTTGGAATT
ATAAAATATAGGTAAGTACGTAATTAAATATTTTG

#### Synthetic Fas Peptides.

Extracellular:

44HLP CPP GER KAR D56

Transmembrane:

157NLG WLC LLL LPI PLI V172

Truncated (-TM):

 $^{147} T \ K \ C \ K \ E \ E^{152} \ ^{174} V \ K \ R \ K \ E \ V^{179}$ 

Numbered according to full length fas polypeptide sequence.

FIGURE 4

8/11
FIGURE 5
ANTI-FAS INDUCED DEATH OF WIL-2 CELLS TRANSFORMED WITH Fas-ΔTM
No Anti-Fas

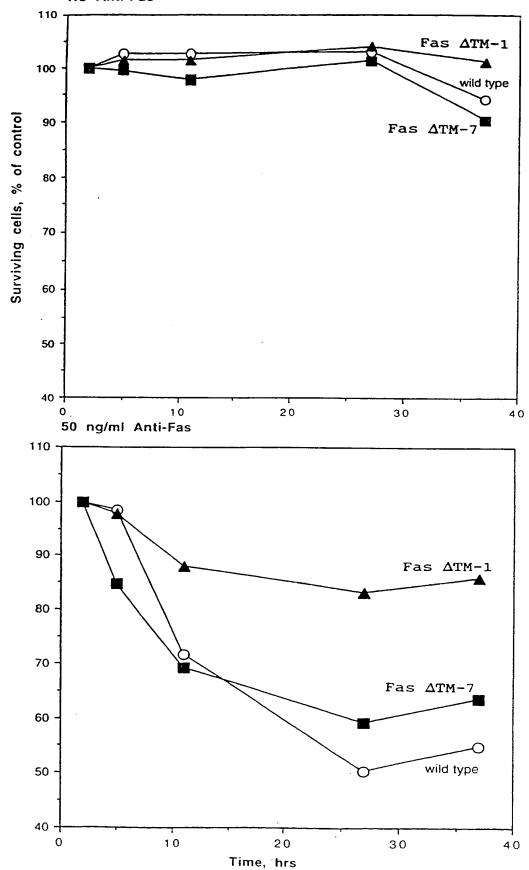
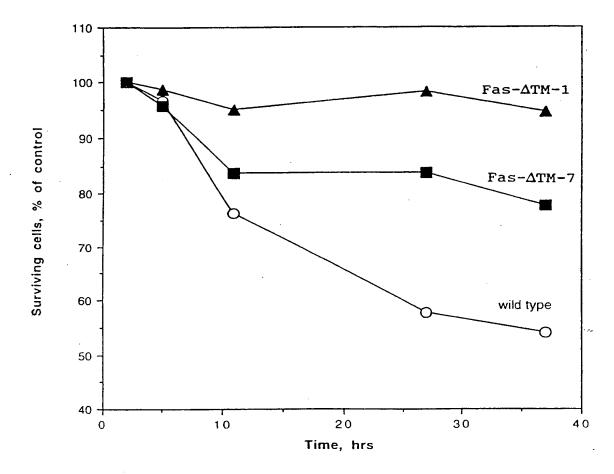


FIGURE 6

# ANTI-FAS INDUCED DEATH OF WIL-2 CELLS TRANSFORMED WITH Fas-ATM (Anti-Fas was added 24 h after seeding)



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#### FIGURE 7

# WIL2/Fas $\Delta$ TM Transformants: Cell Surface Fas

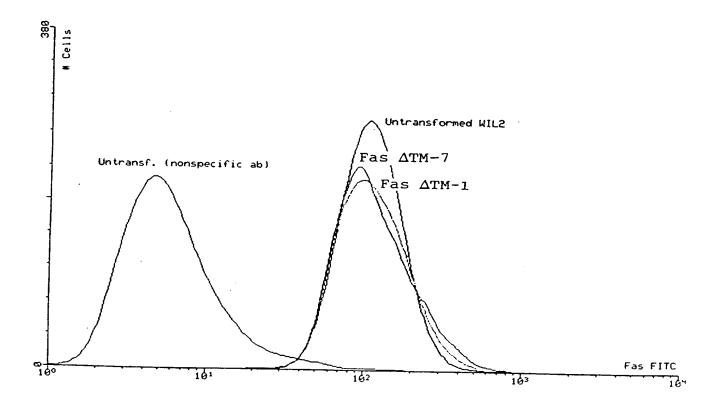
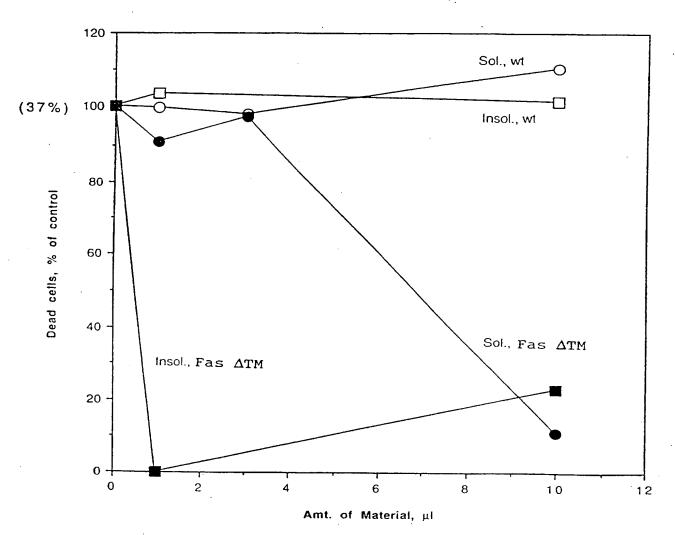


FIGURE 8

# Baculovirus-Fasatm Preparations Prevent Anti-Fas Induced Death of WIL-2 Cells



International application No. PCT/US94/13173

A. CLASSIFICATION OF SUBJECT MATTER  IPC(6): Please See Extra Sheet. US CL: Please See Extra Sheet. According to International Patent Classification (IPC) or to both national classification and IPC  B. FIELDS SEARCHED  Minimum documentation searched (classification system followed by classification symbols)  U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched  Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
US CL: Please See Extra Sheet. According to International Patent Classification (IPC) or to both national classification and IPC  B. FIELDS SEARCHED  Minimum documentation searched (classification system followed by classification symbols)  U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
According to International Patent Classification (IPC) or to both national classification and IPC  B. FIELDS SEARCHED  Minimum documentation searched (classification system followed by classification symbols)  U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
B. FIELDS SEARCHED  Minimum documentation searched (classification system followed by classification symbols)  U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
U.S.: 435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2  Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
Please See Extra Sheet.			
C. DOCUMENTS CONSIDERED TO BE RELEVANT			
Category* Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No			
Y THE JOURNAL OF EXPERIMENTAL MEDICINE, Volume 178, 1-28			
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issued August 1993, J. Wu et al., "Autoimmune disease in mice due to integration of an endogenous retrovirus in an			
apoptosis gene", pages 461-468, see entire document.			
apoptosis gene , pages 401-408, see entire document.			
Y THE JOURNAL OF EXPERIMENTAL MEDICINE, Volume 177, 1-28			
issued January 1993, E. Rouvier et al., "Fas involvement in			
Ca <sup>2+</sup> -independent T cell-mediated cytotoxicity", pages 195-			
200, see entire document.			
200, 000 Cittle document.			
X Further documents are listed in the continuation of Box C. See patent family annex.			
Special categories of cited documents: "T" later document published after the international filing date or priority			
"A" document defining the general state of the art which is not considered to be of particular relevance date and not in conflict with the application but cited to understand the principle or theory underlying the invention			
E* earlier document published on or after the international filing date  *X*  document of particular relevance; the claimed invention cannot be considered to involve an inventive step			
L document which may throw doubts on priority, claim(s) or which is when the document is taken alone cited to establish the publication date of another citation or other			
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means being obvious to a person skilled in the art			
document published prior to the international filing date but later than the priority date claimed document member of the same patent family			
Date of the actual completion of the international search  Date of mailing of the international search report			
21 FEBRUARY 1995 0 3 MAR 1995			
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 2021 BRIAN R. STANTON			
Vasimigion, 5.C. 2021			
Facsimile No. (703) 305-3230 Telephone No. (703) 308-0196 orm PCT/ISA/210 (second sheet)(July 1992)★			

International application No. PCT/US94/13173

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C (C :	W. DOGULARIAM CONTROL	
	ation). DOCUMENTS CONSIDERED TO BE RELEVANT	· · · · · · · · · · · · · · · · · · ·
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim N
Y	Kramer et al., "APOPTOSIS: THE MOLECULAR BASIS OF CELL DEATH", published 1991 by Cold Spring Harbor Laboratory Press (N.Y.), pages 87-99, see entire chapter.	1-28
Y	NATURE, Volume 364, issued 26 August 1993, J. Ogasawara et al., "Lethal effects of the anti-FAS antibody in mice", pages 806-809, see entire document.	1-28
ď	SCIENCE, Volume 261, issued 09 July 1993, N. Zhu et al., "Systemic gene expression after intravenous DNA delivery in adulmice", pages 209-211, see entire document.	t 1-28
	JOURNAL OF BIOLOGICAL CHEMISTRY, Volume 268, Number 22, issued 05 August 1993, K. Hsu et al., "Differential expression and ligand binding properties of tumor necrosis factor receptor chimeric mutants", pages 16430-16436, see entire document.	1-28
	CELL, Volume 75, issued 17 December 1993, T. Suda et al., "Molecular cloning and expression of the fas ligand, a novel member or the tumor necrosis factor family", pages 1169-1178, see entire document.	1-28
1   1   f   r	EUROPEAN JOURNAL OF IMMUNOLOGY, Volume 22, issued 1992, W.A.M. Loenen et al., "The CD27 membrane receptor, a symphocyte-specific member of the nerve growth factor receptor family, gives rise to a soluble form by protein processing that does not involve receptor endocytosis", pages 447-456, see entire document.	

Form PCT/ISA/210 (continuation of second sheet)(July 1992)★

International application No. PCT/US94/13173

Box 1 Observations where certain claims were f und unsearchable (Continuation of item 1 f first sheet)
This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
2. Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
Picase See Extra Sheet.
1. X As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark on Protest  The additional search fees were accompanied by the applicant's protest.  No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet(1))(July 1992)\*

International application No. PCT/US94/13173

## A. CLASSIFICATION OF SUBJECT MATTER: IPC (6):

A01N 43/04; A61K 31/70, 35/16, 39/00; C07H 17/00; C07K 14/00, 14/435, 14/47, 14/705, 16/00; C12N 1/00, 5/06, 15/00, 15/09, 15/12, 15/87; G01N 33/53

## A. CLASSIFICATION OF SUBJECT MATTER: US C:

435/6, 7.1, 7.2, 7.21, 172.1, 172.3, 240.2, 240.21, 320.1; 514/2, 44; 530/350, 387.1; 536/23.1, 23.5, 24.1; 800/2

#### **B. FIELDS SEARCHED**

Electronic data bases consulted (Name of data base and where practicable terms used):

Databases: APS, ANABSTR, AQUASCI, BIOBUSINESS, BIOSIS, BIOTECHDS, CA, CABA, CANCERLIT, CAPREVIEWS, CONFSCI, DISSABS, DRUGB, DRUGLAUNCH, DRUGNL, DURGU, EMBASE, FSTA, GENBANK, HEALSAFE, IFIPAT, JICST-E, JPNEWS, LIFESCIE, MEDLINE

Search Terms: Barr?/au; shaprio?/au; keifer?/au; fas; tm; transmembran?; membran?; solubl?; secret?; antibod?; transgen?; per; polymerase; chain; reaction; gene; therap?

# BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be examined, the appropriate additional examination fees must be paid.

Group I, claims 1-7, drawn to DNA, vectors and host cells comprising a gene encoding FasΔTM.

Group II, claim 8, drawn to transgenic animals comprising a gene for FasΔTM.

Group III, claims 9-13, 16 and 23, drawn to Fas \( \Dag{TM} \) proteins and methods of using said proteins.

Group IV, claims 14, 15 and 17-19, drawn to antibodies that recognize FasΔTM and methods of using said antibodies.

Group V, claims 20-22, drawn to nucleic acids detection assays.

Group VI, claims 24-28, drawn to gene therapy methods using a gene for FasΔTM.

The inventions listed as Groups I-VI do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

The inventions of groups I-IV are distinct, one from the other because they are drawn to materially different compositions that require divergent areas of search, consideration and analysis and that may be used in materially different fashions. For example, the DNA of the invention of group I may be used to prepare recombinant proteins or for use as hybridization probes, which does not involve the in vivo considerations required for analysis of the genetic alteration of an animal, such as those of the invention of group II. The DNA of group I also represents a materially different composition than that of the proteins of group III (comprised of amino acids rather than nucleic acids) and comprises a distinct activity than that of an antibody (group IV) which is a specific, protein based binding reagent). Thus, each of the compositions of the inventions of groups I-IV represent distinct chemical compounds with divergent uses that require disparate considerations based upon their chemical and biological properties.

The inventions of group I and either of the inventions of groups V or VI are distinct, one from the other because the DNA of group I may be used in materially different fashions as evidenced by its use in the detection assays of group V and the therapeutic application of the invention of group VI.

The inventions of groups II-IV and either of the inventions of groups V or VI are distinct, one from the other because the former groups of compositions comprise proteins which are not utilized in the methods of groups V and VI that are based upon nucleic acids.

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The methods of groups V and VI are distinct, one from the other because they are drawn to materially different methods that require divergent areas of search and consideration. In the case of the invention of group VI, the methods are directed to in vivo gene therapy which require consideration of means of administration, targeting and expression and exogenous nucleic acids while the methods of group V are predicated upon in vitro diagnostics which do not include in vivo considerations.

For the reasons stated above, the listed groups of inventions are not so linked by any special technical feature within the meaning of PCT Rule 13.2 so as to comprise a single inventive concept.

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